

- <110> Creech, Christopher D.
 Jegla, Timothy J.
 ICAgen, Inc.
- <120> CNG3B: A Novel Cyclic Nucleotide-Gated Cation Channel
- <130> 018512-006010US
- <140> US 09/855,828
- <141> 2001-05-14
- <150> US 60/204,445
- <151> 2000-05-15
- <160> 16
- <170> PatentIn Ver. 2.1
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- Glu Glu Lys Ser Leu Lys Thr Lys Ser Thr Pro Val Thr Ser Glu Glu 50 55 60
- Pro His Thr Asn Ile Gln Asp Lys Leu Ser Lys Lys Asn Ser Ser Gly 65 70 75 80
- Asp Leu Thr Thr Asn Pro Asp Pro Gln Asn Ala Ala Glu Pro Thr Gly 85 90 95
- Thr Val Pro Glu Gln Lys Glu Met Asp Pro Gly Lys Glu Gly Pro Asn 100 105 110
- Ser Pro Gln Asn Lys Pro Pro Ala Ala Pro Val Ile Asn Glu Tyr Ala 115 120 125
- Asp Ala Gln Leu His Asn Leu Val Lys Arg Met Arg Gln Arg Thr Ala 130 135 140
- Leu Tyr Lys Lys Leu Val Glu Gly Asp Leu Ser Ser Pro Glu Ala 145 150 155 160

- Ser Pro Gln Thr Ala Lys Pro Thr Ala Val Pro Pro Val Lys Glu Ser 165 170 175
- Asp Asp Lys Pro Thr Glu His Tyr Tyr Arg Leu Leu Trp Phe Lys Val
- Lys Lys Met Pro Leu Thr Glu Tyr Leu Lys Arg Ile Lys Leu Pro Asn 195 200 205
- Ser Ile Asp Ser Tyr Thr Asp Arg Leu Tyr Leu Leu Trp Leu Leu Leu 210 215 220
- Val Thr Leu Ala Tyr Asn Trp Asn Cys Trp Phe Ile Pro Leu Arg Leu 225 230 235 240
- Val Phe Pro Tyr Gln Thr Ala Asp Asn Ile His Tyr Trp Leu Ile Ala 245 250 255
- Asp Ile Ile Cys Asp Ile Ile Tyr Leu Tyr Asp Met Leu Phe Ile Gln 260 265 270
- Pro Arg Leu Gln Phe Val Arg Gly Gly Asp Ile Ile Val Asp Ser Asn 275 280 285
- Glu Leu Arg Lys His Tyr Arg Thr Ser Thr Lys Phe Gln Leu Asp Val 290 295 300
- Ala Ser Ile Ile Pro Phe Asp Ile Cys Tyr Leu Phe Phe Gly Phe Asn 305 310 315 320
- Pro Met Phe Arg Ala Asn Arg Met Leu Lys Tyr Thr Ser Phe Phe Glu 325 330 335
- Phe Asn His His Leu Glu Ser Ile Met Asp Lys Ala Tyr Ile Tyr Arg 340 345 350
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- Cys Val Tyr Tyr Trp Ala Ser Asn Tyr Glu Gly Ile Gly Thr Thr Arg 370 375 380
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Asp Leu Thr Thr Asn Pro Asp Pro Gln Asn Ala Ala Glu Pro Thr Gly
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Thr Val Pro Glu Gln Lys Glu Met Asp Pro Gly Lys Glu Gly Pro Asn 100 105 110

Ser Pro Gln Asn Lys Pro Pro Ala Ala Pro Val Ile Asn Glu Tyr Ala 115 120 125

Asp Ala Gln Leu His Asn Leu Val Lys Arg Met Arg Gln Arg Thr Ala 130 135 140

Leu Tyr Lys Lys Lys Leu Val Glu Gly Asp Leu Ser Ser Pro Glu Ala 145 150 155 160

Ser Pro Gln Thr Ala Lys Pro Thr Ala Val Pro Pro Val Lys Glu Ser 165 170 175

Asp Asp Lys Pro Thr Glu His Tyr Tyr Arg Leu Leu Trp Phe Lys Val 180 185 190

Lys Lys Met Pro Leu Thr Glu Tyr Leu Lys Arg Ile Lys Leu Pro Asn 195 200 205

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Val Thr Leu Ala Tyr Asn Trp Asn Cys Trp Phe Ile Pro Leu Arg Leu 225 230 235 240

Val Phe Pro Tyr Gln Thr Ala Asp Asn Ile His Tyr Trp Leu Ile Ala 245 250 255

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Gly Thr Lys Val Leu Val Thr Leu Lys Ala Gly Ser Val Phe Gly Glu

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Gln Glu Ile Leu Val His Tyr Pro Asp Ser Glu Arg Ile Leu Met Lys 625 630 635 640

Lys Ala Arg Val Leu Leu Lys Gln Lys Ala Lys Thr Ala Glu Ala Thr 645 650 655

Pro Pro Arg Lys Asp Leu Ala Leu Leu Phe Pro Pro Lys Glu Glu Thr 660 665 670

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Arg Pro Glu Cys Thr Ala Ser Pro Ile Ala Val Glu Glu Pro His 755 760 765

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- Gly Ile Ala Met Glu Thr Arg Gly Leu Ala Asp Ser Gly Gln Gly Ser 50 55 60
- Phe Thr Gly Gln Gly Ile Ala Arg Leu Ser Arg Leu Ile Phe Leu Leu 65 70 75 80
- Arg Arg Trp Ala Ala Arg His Val His Gln Asp Gln Gly Pro Asp 85 90 95
- Ser Phe Pro Asp Arg Phe Arg Gly Ala Glu Leu Lys Glu Val Ser Ser 100 105 110
- Gln Glu Ser Asn Ala Gln Ala Asn Val Gly Ser Gln Glu Pro Ala Asp 115 120 / 125
- Arg Gly Arg Ser Ala Trp Pro Leu Ala Lys Cys Asn Thr Asn Thr Ser 130 135 140
- Asn Asn Thr Glu Glu Glu Lys Lys Thr Lys Lys Asp Ala Ile Val
- Val Asp Pro Ser Ser Asn Leu Tyr Tyr Arg Trp Leu Thr Ala Ile Ala 165 170 175
- Leu Pro Val Phe Tyr Asn Trp Tyr Leu Leu Ile Cys Arg Ala Cys Phe 180 185 190
- Asp Glu Leu Gln Ser Glu Tyr Leu Met Leu Trp Leu Val Leu Asp Tyr 195 200 205
- Ser Ala Asp Val Leu Tyr Val Leu Asp Val Leu Val Arg Ala Arg Thr 210 215 220
- Gly Phe Leu Glu Gln Gly Leu Met Val Ser Asp Thr Asn Arg Leu Trp 225 230 235 240
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- Tyr Phe Ala Ile Ser Lys Phe Ile Gly Phe Gly Thr Asp Ser Trp Val 325 330 335
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Ala Ser Thr Ser Glu Glu Ser Glu Asn Glu Asn Pro His Ala Arg Gly
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Gln Tyr Leu Pro Gly Ala Ile Ala Leu Phe Asn Val Asn Asn Ser Ser 85 90 95

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Lys Ser Lys Ser Asp Asp Lys Asn Glu Asn Lys Asn Asp Pro Glu Lys 115 120 125

Lys Lys Lys Lys Lys Asp Lys Glu Lys Lys Lys Glu Glu Lys Ser 130 135 140

Lys Asp Lys Lys Glu Glu Glu Lys Lys Glu Val Val Ile Asp Pro 145 150 155 160

Ser Gly Asn Thr Tyr Tyr Asn Trp Leu Phe Cys Ile Thr Leu Pro Val 165 170 175

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